! Sequence: momp ! Sequence: incC ! Sequence: pomp91a

CLUSTAL W (1.83) multiple sequence alignment

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momp
incC
            MKQMRLWGFLFLSSFCQVSYLRANDVLLPLSGIHSGEDLELFTLRSSSPTKTTYSLRKDF
pomp9la
                        -----MKKLLKSVLVFAALSSASSLQALPVG--
momp
                     ______MTSPIPFQ--
incC
             IVCDFAGNSIHKPGAAFLNLKGDLFFINSTPLAALTFKNIHLGARGAGLFSESNVTFKGL
pomp9la'
             momp
             -----SSGDASFLAEQPQQLPSTSESQLVTQLLTMMKHTQALSETVLQQ
incC
             HSLVLENNESWGGVLTTSGDLSFINNTSVLCQNNISYGFGGALLLQGRKSKALFFRDNRG
pomp91a
                        . ...: *:: :
            DRVLKTDVNKEFQMGAKPTTDTGNSAAPS-----
πφmφ
             QRDRLPTASIILQVGGAFTGGAGAPFQFG-----
incC
            TILFLKNKAVNQDESHPGYGGAVSSISPGSPITFADNQEILFQENEGELGGAIYNDQGAI
pomp91a
                            .: . *.
              qmom
             _____PADDHHHPIPPPVVPAQIETEITTIRSELQLMR--
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qmom
             --- STLQQSTKGARTGVLVVTAILMTISLLAIIIILAVLGFTG---VLPQVALLMQGETN
incC
            KGSIVFEENSATAGGAIAVNAVCDINAQGPVRFINNSALGLNGGAIYMQATGSILRLHAN
pomp91a
                                 .
             _____FAWSVGARAALWECGCATLG----
momp
incC
            QGDIEFCGNKVRSQFHSHINSTSNFTNNAITIQGAPREF$LSANEGHRICFYDPIISATE
pomp91a
incC
            NYNSLYINHQRLLEAGGAVIFSGARLSPEHKKENKNKTSIINQPVRLCSGVLSIEGGA1L
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                    ._____ASFQYAQSKPKVEELNVLCNAAEF
momp
             _____TLGLILTNKNTPLPAS-----
incC
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pomp91a
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momp
incC ..
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qmom
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cmom
incC
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pomp91a	VNNNYLNNSEVIPLOHLCVFGGPVYQIMEQNPKQSSNNLLVQHAGHNVGARIPFSFNTIL
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momp incC pomp91a	GTSRGSWRNYGWSGSVGMSYAYPKGIRYLKMTPFVDLQYTKLVQNPFVETGYDPRYFSSS
momp incC pomp91a	EMTNLSLPIGIALEMRFIGSRSSLFLQVSTSYIKDLRRVNPQSSASLVLNHYTWDIQGVP
momp incC pomp91a	LGKEALNITLNSTIKYKIVTAYMGISSTQREGSNLSANAHAGLSLSF

[&]quot;*" means that the residues or nucleotides in that column are identical in all sequences in the alignment
"" means that conserved substitutions have been observed

[&]quot;." means that semi-conserved substitutions are observed

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6693087pomp91a seq
947 aa
                                                                                                                                                        PAT 20-FEB-2004
                                                                                                                                   linear
                            AAS37561
LOCUS
                            Sequence 3 from patent US 6693087.
DEFINITION
ACCESSION
                            AAS37561
                            AAS37561.1 GI:42715796
VERSION
DBSOURCE
                            accession AAS37561.1
KEYWORDS
                            Unknown.
SOURCE
                            Unknown.
Unclassified.
     ORGANISM
                                  (residues 1 to 947)
                            Murdin,A.D., Dunn,P.L. and Oomen,R.P.
Nucleic acid molecules encoding POMP91A protein of Chlamydia
Patent: US 6693087-A 3 17-FEB-2004;
Aventis Pasteur Limited; Toronto;
REFERENCE
    AUTHORS
     TTTLE
     JOURNAL
                            CAMBIA Patent Lens: US 6693087
Location/Qualifiers
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ORIGIN
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6686339incc seq
                                                                                                               PAT 20-FEB-2004
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DEFINITION Sequence 3 from patent US 6686339.
                    AAS33023
AAS33023.1 GI:42707452
ACCESSION
VERSION
                     accession AAS33023.1
DBSOURCE
KEYWORDS
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1 (residues 1 to 203)

Murdin,A.D., Dunn,P.L. and Oomen,R.P.

Nucleic acid molecules encoding inclusion membrane protein C of Chlamydia

Patent: Un 6500000 1 10
                     Unknown.
SOURCE
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REFERENCE
   AUTHORS
   TITLE
                    Patent: US 6686339-A 3 03-FEB-2004;
Aventis Pasteur Limited; Toronto;
   JOURNAL
                    CAMBIA Patent Lens: US 6686339
Location/Qualifiers
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181 cfialigtlg liltnkntpl pas
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stephens momp
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                                                                                                                                                               linear
                                   Q46409
                                                                                                                                                                                        BCT 02-MAY-2006
 LOCUS
                                   Major outer membrane protein, serovar D precursor (MOMP).
 DEFINITION
                                    Q46409
 ACCESSION
 VERSION
                                    Q46409
                                                         GI:6707730
                                  swissprot: 10043 Class: standard. created: May 30, 2000. sequence updated: Nov 1, 1996. annotation updated: May 2, 2006. vrefs: x62918.1, CAA44701.1, AF063195.2, AAC31436.2, AE001273.1, AP063195.2 AAC31436.2, AE001273.1, AP063195.2 AAC31436.2, AE001273.1, AP063195.2 AAC31436.2 AE001273.1, AP063195.2 AE001273.1, AE001273.
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GenomeReviews:AE001273_GR, InterPro:IPR000604, Pfam:PF01308,
                                   PRINTS: PRO1334
                                 Complete proteome; Ion transport; Membrane; Outer membrane; Porin; Signal; Transmembrane; Transport. Chlamydia trachomatis Chlamydia trachomatis Bacteria; Chlamydiae; Chlamydiales; Chlamydiae; Chlamydiae; Chlamydiae; Chlamydiae; Chlamydiae; Chlamydiae; Chlamydiae.
KEYWORDS
SOURCE '
      ORGANISM
REFERENCE
                                  Sayada, C., Denamur, E. and Elion, J.
Complete sequence of the major outer membrane protein-encoding gene
of Chlamydia trachomatis servovar Da
      AUTHORS
      TITLE
                                   Gene 120 (1), 129-130 (1992)
1398119
      JOURNAL
        PUBMED
                                  NUCLEOTIDE SEQUENCE [GENOMIC DNA].
STRAIN=D/B-120
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      REMARK
REFERENCE
                                  Stothard, D.R., Boguslawski, G. and Jones, R.B. Phylogenetic analysis of the Chlamydia trachomatis major outer
     AUTHORS
      TITLE
                                  membrane protein and examination of potential pathogenic
                                   determinants
                                   Infect. Immun. 66 (8), 3618-3625 (1998)
     JOURNAL
                                   9673241
        PUBMED
                                  NUCLEOTIDE SEQUENCE [GENOMIC DNA].
     REMARK
                                  STRAIN-D/IU-71960
                                 3 (residues 1 to 393)
Stephens,R.S., Kalman,S., Lammel,C., Fan,J., Marathe,R.,
Aravind,L., Mitchell,W., Olinger,L., Tatusov,R.L., Zhao,Q.,
Koonin,E.V. and Davis,R.W.
REFERENCE
     AUTHORS
                                  Genome sequence of an obligate intracellular pathogen of humans:
     TITLE
                                  Chlamydia trachomatis
Science 282 (5389), 754-759 (1998)
     JOURNAL
        PUBMED
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On Sep 27, 2005 this sequence version replaced gi:7442973.

[FUNCTION] Structural rigidity of the outer membrane of elementary bodies and porin forming, permitting diffusion of solutes through the intracellular reticulate body membrane.

[SUBUNIT] Disulfide bond interactions within and between MOMP
COMMENT
                               -molecules and other components form high molecular-weight
                              .,[SUBCELLULAR LOCATION] Bacterial cell outer membrane; multi-pass
                                 membrane protein.
                                  [SIMILARITY] Belongs to the chlamydial OMP family.
                                                            Location/Qualifiers
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                                                           /organism="Chlamydia trachomatis"
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                                                                                                              Page 1
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